

1102-98.TXT SEQUENCE LISTING

<110> Kainoh, Mie Tanaka, Toshiaki

<120> Chimeric Proteins, their Heterodimer Complexes, and Platelet Substitutes

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aat Asn	gtt Val	acc Thr	tgc Cys 1060	Trp	ttg Leu	aaa Lys	gac Asp	gtt Val 106!	His	atg Met	aaa Lys	gga Gly	gaa Glu 1070	Tyr	ttt Phe	3216
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cct gag ata t Pro Glu Ile T 1105	at gtg att yr Val Ile 1110	Glu Asp .	Asn Thr	gtt acg a Val Thr I 1115	tt ccc ctg le Pro Leu	atg 3360 Met 1120
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ccc aaa tct t Pro Lys Ser C 1		Thr His				3807
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gcc ctc cca g Ala Leu Pro A 1		Glū Lys				4255
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1102-98.TXT Gly Thr Ser Asp Val Ile Thr Gly Ser Ile Gln Val Ser Ser Arg Glu Ala Asn Cys Arg Thr His Gln Ala Phe Met Arg Lys Asp Val Arg Asp 565 570 575 Ile Leu Thr Pro Ile Gln Ile Glu Ala Ala Tyr His Leu Gly Pro His
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1102-98.TXT Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His
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1102-98.TXT Leu Thr Asn Lys Gly Glu Val Phe Asn Glu Leu Val Gly Lys Gln Arg lle Ser Gly Asn Leu Asp Ser Pro Glu Gly Gly Phe Asp Ala Ile Met 245 250 255 Gln Val Ala Val Cys Gly Ser Leu Ile Gly Trp Arg Asn Val Thr Arg 260 265 270 Leu Leu Val Phe Ser Thr Asp Ala Gly Phe His Phe Ala Gly Asp Gly 275 280 Lys Leu Gly Gly Ile Val Leu Pro Asn Asp Gly Gln Cys His Leu Glu 295 300 Asn Asn Met Tyr Thr Met Ser His Tyr Tyr Asp Tyr Pro Ser Ile Ala 305 315 310 His Leu Val Gln Lys Leu Ser Glu Asn Asn Ile Gln Thr Ile Phe Ala 325 330 335 Val Thr Glu Glu Phe Gln Pro Val Tyr Lys Glu Leu Lys Asn Leu Ile 340 345 _ 350 _ _ Pro Lys Ser Ala Val Gly Thr Leu Ser Ala Asn Ser Ser Asn Val Ile 360 Gln Leu Ile Ile Asp Ala Tyr Asn Ser Leu Ser Ser Glu Val Ile Leu 370 375 380 Glu Asn Gly Lys Leu Ser Glu Gly Val Thr Ile Ser Tyr Lys Ser 385 390 395 Cys Lys Asn Gly Val Asn Gly Thr Gly Glu Asn Gly Arg Lys Cys Ser Asn Ile Ser Ile Gly Asp Glu Val Gln Phe Glu Ile Ser Ile Thr Ser 420 425 Asn Lys Cys Pro Lys Lys Asp Ser Asp Ser Phe Lys Ile Arg Pro Leu 435 440 445 Gly Phe Thr Glu Glu Val Glu Val Ile Leu Gln Tyr Ile Cys Glu Cys 450 455 460 Glu Cys Gln Ser Glu Gly Ile Pro Glu Ser Pro Lys Cys His Glu Gly 470 475 Asn Gly Thr Phe Glu Cys Gly Ala Cys Arg Cys Asn Glu Gly Arg Val 485 490 495 Gly Arg His Cys Glu Cys Ser Thr Asp Glu Val Asn Ser Glu Asp Met 500 510 Asp Ala Tyr Cys Arg Lys Glu Asn Ser Ser Glu Ile Cys Ser Asn Asn 515 520 525 Gly Glu Cys Val Cys Gly Gln Cys Val Cys Arg Lys Arg Asp Asn Thr 530 _ 535 540 Asn Glu Ile Tyr Ser Gly Lys Phe Cys Glu Cys Asp Asn Phe Asn Cys 545 550 555 560 Asp Arg Ser Asn Gly Leu Ile Cys Gly Gly Asn Gly Val Cys Lys Cys 565 570 _ 575 Arg Val Cys Glu Cys Asn Pro Asn Tyr Thr Gly Ser Ala Cys Asp Cys 580 585 _ 590 Ser Leu Asp Thr Ser Thr Cys Glu Ala Ser Asn Gly Gln Ile Cys Asn 595 600 Gly Arg Gly Ile Cys Glu Cys Gly Val Cys Lys Cys Thr Asp Pro Lys 610 620 Phe Gln Gly Gln Thr Cys Glu Met Cys Gln Thr Cys Leu Gly Val Cys 625 635 635 640 Ala Glu His Lys Glu Cys Val Gln Cys Arg Ala Phe Asn Lys Gly Glu 65Ŏ 645 Lys Lys Asp Thr Cys Thr Gln Glu Cys Ser Tyr Phe Asn Ile Thr Lys 660 665 670 670 Val Glu Ser Arg Asp Lys Leu Pro Gln Pro Val Gln Pro Asp Pro Val 675 680 685 Ser His Cys Lys Glu Lys Asp Val Asp Asp Cys Trp Phe Tyr Phe Thr 695 700 Tyr Ser Val Asn Gly Asn Asn Glu Val Met Val His Val Val Glu Asn 705 710 715 720 Pro Glu Cys Pro Thr Gly Pro Glu Asp Pro Glu Glu Pro Lys Ser Cys Page 26

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1102-98.TXT
1180

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1205
1210

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Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser
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Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser
1365
Leu Ser Leu Ser Pro Gly Lys
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